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(54) Recombinant gamma interferons, DNA therefor and pharmaceutical compositions containing them.

(57) The present invention relates to novel, stable recombinant gamma interferons exhibiting in greater or less degree the antiviral and antiproliferative activity in humans and pH 2 labile properties characteristic of native human gamma interferon. The amino acid sequence of such an interferon comprises, from the N terminus, at least

X - Y - ASP . . . . . THR  
1 2 126

wherein X is a methionine residue or hydrogen and Y is a glutamine residue or, where X is hydrogen, Y is either a glutamine or a pyroglutamate residue.

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RECOMBINANT GAMMA INTERFERONS, DNA  
THEREFOR AND PHARMACEUTICAL COMPOSITIONS  
CONTAINING THEM

5

The present invention relates to the field of recombinant DNA technology, to means and methods utilizing such technology in the preparation of recombinant gamma interferons having enhanced stability, to their production  
10 and to the various products of such production and their uses.

This application is related to GB 2107718.

The publications and other materials used herein to illuminate the background of the invention and in  
15 particular cases to provide additional details respecting its practice are for convenience numerically referenced in the following text and respectively grouped in the appended bibliography.

Human interferons can be classified in three groups  
20 on the basis of different antigenicity and biological and biochemical properties. The first group comprises a family of leukocyte interferons which are normally produced mainly by constituent cells

of human blood upon viral induction. These have been microbially produced and found to be biologically active (1,2,3). Their biological properties have prompted their use in the clinic as therapeutic agents for the treatment of viral infections and malignant conditions (4).

In the second group is human fibroblast interferon, normally produced by fibroblasts upon viral induction, which has likewise been microbially produced and found to exhibit a wide range of biological activities (5). Clinical trials also indicate its potential therapeutical value. The leukocyte and fibroblast interferons exhibit very clear similarities in their biological properties despite the fact that the degree of homology at the amino acid level is relatively low. Both groups of interferons contain from about 165 to about 166 amino acids and are acid stable proteins.

Human gamma interferon (also variously referred to as immune interferon,  $\gamma$ -interferon, IIF or IFN- $\gamma$ ) exhibits the antiviral and anti-proliferative properties characteristic of the interferons but, in contrast to leukocyte and fibroblast interferons, is pH 2 labile. Prior to the production of gamma interferons via recombinant DNA technology, it had been produced mainly upon mitogenic induction of lymphocytes. Human gamma interferon is clearly antigenically distinct from the leukocyte and fibroblast interferons. Gray, Goeddel and co-workers were the first to report expression of a recombinant gamma interferon (6), which has proven to exhibit the characteristic properties of human gamma interferon, i.e., anti-viral and anti-proliferative activity coupled with pH 2 lability. The recombinant gamma interferon of Gray and Goeddel, as produced in *E. coli*, consisted of 146 amino acids, the N-terminal portion of the molecule commencing with the sequence CYS-TYR-CYS-. Derynck and others subsequently reported (7) a further recombinant gamma interferon having the same N-terminus and a single amino acid substitution, the polypeptide perhaps constituting an allelic variant of that earlier reported in reference (6). Other workers have reported the production of still further recombinant gamma interferons in which one or more of the amino acids present in

Goeddel and Gray's original publication (6) have allegedly been substituted.

For example, Alton et al. (17) report on a series of IFN-gammas wherein a single amino acid substitution at position 81 of the Gray et al. (6) gamma interferon resulted in an IFN-gamma that retained only 70 percent of the activity (on a relative basis) and wherein an additional deletion of the cys-tyr-cys at positions 1, 2, 3 of this IFN-gamma further reduced relative activity resulting in an IFN-gamma having only 49 percent of the Gray et al. (6) gamma interferon.

In our hands, recombinant gamma interferons whose N-terminal amino acid sequence comprises cysteine residues have proven problematic from the standpoint of oligomerization which may involve participation of sulfhydryl groups of one or more of the cysteine residues in disulfide bond formation. Our inability to completely reduce these putative disulfide linkages suggests the problem may be more complex, possibly also involving reaction through the hydroxyl function of the cysteine-bounded tyrosine residue. These recombinant interferons have proven somewhat unstable and, whether resulting from such instability or otherwise, have proven of less than optimal utility.

The present invention relates to recombinant gamma interferons having enhanced stability and activity. In the most preferred embodiment we provide a recombinant gamma interferon comprising the amino acid sequence (hereinafter "full sequence"), extending from the N-terminus:

X-Y-ASP-PRO-TYR-VAL-LYS-GLU-ALA-GLU-ASN-LEU-LYS-LYS-TYR-PHE-ASN-ALA-  
 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17  
 GLY-HIS-SER-ASP-VAL-ALA-ASP-ASN-GLY-THR-LEU-PHE-LEU-GLY-ILE-LEU-LYS-  
 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34  
 ASN-TRP-LYS-GLU-GLU-SER-ASP-ARG-LYS-ILE-MET-GLN-SER-GLN-ILE-VAL-SER-  
 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51  
 PHE-TYR-PHE-LYS-LEU-PHE-LYS-ASN-PHE-LYS-ASP-ASP-GLN-SER-ILE-GLN-LYS-  
 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68

SER-VAL-GLU-THR-ILE-LYS-GLU-ASP-MET-ASN-VAL-LYS-PHE-PHE-ASN-SER-ASN-  
69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85

LYS-LYS-LYS-ARG-ASP-ASP-PHE-GLU-LYS-LEU-THR-ASN-TYR-SER-VAL-THR-ASP-  
86 87 88 89 90 91 92 93 94 95 96 97 98 99 100 101 102

5 LEU-ASN-VAL-GLN-ARG-LYS-ALA-ILE-HIS-GLU-LEU-ILE-GLN-VAL-MET-ALA-GLU-  
103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118 119

LEU-SER-PRO-ALA-ALA-LYS-THR-GLY-LYS-ARG-LYS-ARG-SER-GLN-MET-LEU-PHE-  
120 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136

10 ARG-GLY-ARG-ARG-ALA-SER-GLN  
137 138 139 140 141 142 143

wherein X is a methionine residue or hydrogen and Y is a glutamine residue or, where X is a hydrogen, either a glutamine or pyroglutamate residue, as well as various analogs thereof in which specified portions of the carboxy terminal of the foregoing sequence are absent. The invention also provides corresponding cloned genes, expression vectors comprising them, and transformants useful in the production through recombinant DNA technology of the interferons of the invention. The preferred recombinant gamma interferons of the invention (as compared to those previously characterized in the literature) appear to most closely approximate the true amino acid sequence of native gamma interferon, the latter of which we have now purified from native sources and fully characterized. Preferred embodiments of the invention exhibit greatly improved stability and activity relative to those previously described in the literature.

25 In the drawings:

The manner in which these and other objects of the invention are attained will be apparent from the detailed description which follows and from the accompanying drawings, in which:

30 Fig. 1 illustrates amino acids 1 through 143 of a recombinant gamma interferon of the present invention and DNA sequence encoding the amino acid sequence preceded by a signal sequence, which DNA sequence is flanked by regions of 5'- and 3'-untranslated DNAs.

35 Fig. 2 illustrates schematically a plasmid coding for direct synthesis of a recombinant gamma interferon of the present invention in E. coli and its preparation.

Fig. 3 records data demonstrating the enhanced stability of gamma interferon prepared according to the present invention.

We have learned that native human gamma interferon (i.e.,  
5 that arising from mitogen induction of human peripheral blood  
lymphocytes and subsequent purification) is a polypeptide which  
lacks the CYS-TYR-CYS- N-terminus assigned by Gray et al. to the  
recombinant gamma interferon whose sequence is depicted in (6).  
Tryptic digests of highly purified native gamma interferon in our  
10 hands included sequences whose amino acid composition generally  
corresponded to that of the N-terminal portion of the Gray et al.  
recombinant gamma interferon of (6), less CYS-TYR-CYS. Amino acid  
sequence analysis from the N-terminus of native gamma interferon  
proved unavailing, giving rise to the inference that the alpha amino  
15 acid at the N-terminus of the molecule was protected. Since the  
first amino acid beyond the second cysteine of Gray et al. (6) for  
which the cDNA coded was GLN (glutamine), we surmised that  
cyclization of the GLN residue had left instead pyroglutamate, such  
that the N-terminus was blocked. Removal of pyroglutamate with  
20 pyroglutamate aminopeptidase left a free alpha amino group  
associated with ASP, the next encoded amino acid, and sequence  
analysis could proceed, permitting the first reported  
characterization of native human gamma interferon.

Appropriate alteration of cDNA for CYS-TYR-CYS- containing  
25 recombinant human gamma interferon permitted the direct expression  
in E. coli of novel recombinant gamma interferon from a cDNA  
encoding the protein whose full sequence is set out supra, X being  
MET and Y being GLN. The N-terminal methionine is an artifact  
encoded by the mRNA translational "start" signal AUG which, in the  
30 particular case of E. coli expression exemplified, is not processed  
away by host systems. In other microbial systems, e.g., pseudomonas,  
methionine may be removed; it does not in any event appear required  
for activity. Where methionine is removed and depending upon the  
system employed, the GLN residue may cyclize to the pyroglutamate  
35 form, again without any believed impairment of activity.

In our hands, the CYS-TYR-CYS- containing recombinant gamma interferon earlier reported by Gray and co-workers benefited from formulation with human serum albumin in aid of stabilization. The presence of serum albumin in the final lyophilized product, however, requires that certain quality control steps be performed in advance of lyophilization rather than upon finished product. In the case of the novel recombinant gamma interferon of present invention, on the other hand, the material in lyophilized form has proven to be sufficiently stable without the inclusion of serum albumin. Where desired, however, the gamma interferons of the invention may be formulated with pharmaceutically acceptable levels of human serum albumin.

Beyond the foregoing, the CYS-TYR-CYS- lacking recombinant human gamma interferons of the invention appear in cytopathic effect inhibition testing to be markedly more active as antiviral agents than their CYS-TYR-CYS- containing analogs. The activity is conventionally assayed in microtiter plates by inhibition of the cytopathic effect (CPE) of encephalomyocarditis virus on human lung carcinoma cells A549. See (12).

The recombinant gamma interferons of the invention include all those comprising amino acids 1 to about 126 of the full sequence provided above. Gamma interferons variously truncated at the carboxy terminal end relative to the full sequence continue to exhibit the characteristic properties of human gamma interferon, albeit at diminished levels in some cases, so long as amino acids 1 to about 126 are present. Indeed, experiments with the CYS-TYR-CYS- containing analog reported at (7) showed that extraneous sequences could be substituted for the amino acid sequence following amino acid 132 (by the present numbering system) without loss of activity. See, e.g., (8). Preliminary evidence in our hands supports the hypothesis that while amino acids 1 to about 126 (THR) are relatively tightly bound in a three-dimensional configuration we

various portions of the sequence downstream from amino acid 126, but not upstream therefrom. Native gamma interferon moieties in our hands include molecules variously extending through amino acids 127, 128, 129, 130, 132 and 134. We have seen fully active recombinant gamma interferon whose amino acid sequence following methionine consisted variously of amino acids (beyond MET) 1 to about 139 and 1 to about 131, the latter obtained by limited digestion of recombinant gamma interferon with trypsin followed by sequence confirmation. Similar trypsin digested fragments variously ending at about amino acids (beyond MET) 128 and 129 retained activity, albeit substantially diminished. On the other hand, material having 125 amino acids (in addition to N-terminal methionine), the threonine at position 126 and following amino acids having been digested away, exhibited less than 1% the activity of undigested material in CPE inhibition assay.

Recombinant derived gamma interferon, in addition to its bearing an initial methionine when produced in hosts where the methionine is not intracellularly cleaved, appears to exhibit a major species having 139 amino acids (based on the numbering system of Figure 1) and a minor species having 143 amino acids. The composition of the two species contains greater than about 95 percent, most preferably greater than about 97 percent of the species having 139 amino acids. Trypsin digestion under limiting conditions likewise removes various portions of the sequence downstream from about amino acid 126. Recombinant gamma interferon in our hands following such limiting trypsin digestion include species variously extending through amino acids 125, (126 with an initial methionine) 129, 131 and 134. These species have retained activity, albeit substantially diminished, in the zone extending from about amino acid 125 to about 129. Species having at least about 129 amino acids, and especially at least about 131 amino acids, i.e., species having from about 129 to 143 amino acids are essentially functionally fully active.

In view of the foregoing, it will be apparent that the invention comprises not only recombinant gamma interferons



exhibiting the full sequence depicted above, but also those, and mixtures of various of those, in which amino acid 143 is absent or in which the amino acid sequence Z→ amino acid 143 is absent, Z being amino acid 127, 128, 129, 130, 131, 132, 133, 134, 135, 136, 137, 138, 139, 140, 141 or 142. It will likewise be apparent that double stranded DNA sequences encoding recombinant gamma interferons according to the invention comprise not only those encoding the full sequence depicted above, but also those encoding only the various carboxy terminal - truncated analogs just described, in the case of each different truncation the following codon(s) encoding translation stop signal(s).

By reference to "recombinant gamma interferon" herein we intend a polypeptide (whether or not glycosylated by the cell in which it is produced) expressed in a transformant cell from a replicable expression vehicle arising from recombinant DNA technology, the polypeptide exhibiting in greater or less degree the antiviral and antiproliferative activity in humans and pH 2 labile properties characteristic of native human gamma interferon.

The recombinant gamma interferons described herein are believed to form "dimers", defined for the purpose of this disclosure as a combination of two such polypeptides each having at least the 1 to about 126 amino acids (which polypeptides may have the same or a different number of amino acids). The nature of the chemical combining mechanism is not fully understood, but is believed to be other than covalent bonding. This combination into dimers appears to occur spontaneously and is believed to be inevitable in the systems described herein. Thus, when the recombinant gamma interferons of the present invention are administered, they will usually be in the dimerized form.

Furthermore, it is to be understood that here, similarly to the disclosures in the literature relating to differently characterized recombinant gamma interferons (8, 9), amino acid substitutions or additions, particularly single amino acid substitutions and addition or substitution of groups of amino acids upstream or downstream from about amino acid 126 or the C-terminus,

in the recombinant gamma interferons disclosed herein are possible without destroying the interferon activity which they possess. It is believed that it will be obvious to those skilled in the art to make such substitutions or additions without departing from the scope of the present invention. Again, the recombinant gamma  
5 interferons of the present invention include species having modifications and allelic variations of the full sequence (see Figure 1) which exhibit biological activity equivalent to or greater than that of the full sequence.

10 Characteristically, purified recombinant gamma interferons will be essentially free of other proteins of human origin, and are so to be distinguished from the native human gamma interferon compositions heretofore available.

15 The invention includes recombinant gamma interferon compositions which (prior to formulation) are greater than about 95 percent pure, preferably more than about 98 percent pure, which compositions are for that reason likewise distinct from native gamma interferons heretofore available.

20 Embodiments of the invention in which the N-terminal amino acid residue is methionyl are likewise distinct from gamma interferons produced within the human body, and appear also for that reason to be distinct, beyond their deletion of CYS-TYR-CYS-, from those whose sequence is reported by Gray et al. (6).

25 The replicable expression vehicles referred to herein are double-stranded DNA moieties, preferably plasmids, comprising an origin of replication, a promoter or promoter-operator, a sequence encoding a ribosome binding site, a codon for a translation start signal, and in proper reading phase therewith a gene encoding the recombinant gamma interferon of interest, followed by codon(s) for a  
30 translation stop. At the present stage the general techniques and lexicography of recombinant DNA technology are well understood to the art-skilled, who are referred in any event to (11) for background information pertinent to the practice of the present invention, mutatis mutandis, in all its embodiments and legally  
35 cognizable equivalents.

ExampleA. Cloning

Recombinant DNA clones containing gamma interferon cDNA sequences were prepared as described in (6) and in the  
5       aforementioned GB 2107718 with messenger  
RNA from induced human peripheral blood lymphocytes. The DNA  
sequence of clone p67 is shown in Figure 1. A 5' untranslated  
region is followed by 69 nucleotides encoding a precursor or signal  
peptide of 23 amino acids, 429 nucleotides coding for a mature  
10       interferon polypeptide of 143 amino acids, and 587 nucleotides of 3'  
untranslated sequence.

B. Expression1. E.Coli Example

To express high levels of recombinant IFN- $\gamma$  in E. coli, the  
15       initiation of protein synthesis must occur at an ATG codon  
immediately preceding the glutamine codon (amino acid one) of the  
mature polypeptide rather than at the ATG of the signal peptide  
(amino acid S1) (Fig. 1). The procedure followed to express the  
cDNA insert of p67 directly in E. coli is outlined in Fig. 2. The  
20       approach was similar to that used to express in E. coli the cDNA  
insert of Gray et al. (6).

An AvaI restriction site located at codon 2 of the  
presumed mature coding sequence was utilized to remove the signal  
peptide coding region. Two synthetic deoxyoligonucleotides were  
25       designed which restore the codons for amino acids 1 and 2,  
incorporate an ATG translational initiation codon, and create an  
XbaI cohesive terminus. These two oligomers were ligated to the  
remainder of the cDNA insert to construct a 1063 base-pair  
synthetic-natural hybrid gene coding for a polypeptide of 144 amino  
30       acids and bounded by XbaI and PstI sites. This gene was inserted  
into the plasmid pLeIF A25 (10) between the XbaI and PstI sites to  
give the expression plasmid p $\gamma$ -CYC5. E. coli strain W3110 (F<sup>-</sup>,  $\lambda$   
- , protrophic) (ATCC No. 27325) was transformed with this plasmid  
to give the host-vector combination E. coli W#110/p $\gamma$ -CYC5.  
35

## 2. Cell Culture Example

Expression of a gene encoding both the signal peptide and gamma interferon, as indicated in Figure 1, was effected in COS-7 cells (16) in the presence of radioactively labeled cysteine and methionine, confirming the production from the gene of mature gamma interferon whose N-terminal amino acids are as indicated in Figure 1 (unlike the case involving E. coli expression, the expression product of mammalian cell systems like that exemplified here lacks N-terminal methionine).

Confluent monolayers of COS-7 cells in 60 mm petri dishes were transfected in duplicate with DNA using the modified DEAE-dextran procedure. Three days after DNA addition, the media was removed. Each set of plates received 2 mls DMEM supplemented with either 100  $\mu$ Ci  $S^{35}$ -methionine or  $S^{35}$ -cysteine. After 16 hours incubation in the presence of the radiolabeled amino acid, the media was removed and 500  $\mu$ l immunoprecipitated using an anti-gamma-interferon monoclonal antibody or an anti-HBsAg monoclonal antibody as the first antibody and a rabbit anti-mouse IgG antibody (Cappel Inc.) as the second antibody. Reaction with the antibody and the subsequent binding to Staphylococcus A cells (Calbiochem) are as described by Berman, P. et al. (18). The samples were resuspended in SDS-mercaptoethanol and electrophoresed on 10% SDS-PAGE gels. The gel was fixed in 7% acetic acid in ethanol, soaked in Enhance (New England Nuclear) fluor solution, dried down, and exposed for two weeks using Kodak AR5 film and an intensifying screen (Dupont).

Plasmids used in this study were pSVgamma69 (11); pDL RI (19), a hepatitis B virus surface antigen expression vector upon which pSVgamma69 was based; and pDL RIgamma Sau, a polycistronic plasmid containing the 830 bp SAU3a fragment of pSVgamma69 (11) spanning the entire gamma-interferon encoding sequences inserted into the EcoRI site of pDL RI. The latter plasmid produces a transcript containing both the gamma-interferon and the HBsAg coding regions.

Comparison of  $S^{35}$ -Cysteine and  $S^{35}$ -Methionine labeled

proteins which react with either anti-gamma-interferon (A) or anti-HBsAg (B) antibodies showed that no material migrating at either the glycosylated (29,000 MW) or monoglycosylated position (18,000 MW) was specifically immunoprecipitated from  $S^{35}$ -Cys labeled pDL RI-gamma Sau transfected cells using anti-gamma-interferon antibody, in contrast to the  $S^{35}$ -met labeled cells which showed the immunoprecipitation of interferon-gamma.

#### C. Fermentation Production

The production of recombinant Human Interferon - Gamma (rIFN- $\gamma$ ) using E. coli W3110/py- CYC5 is carried out in batches ranging in volume from 10 to 1000 liters. After fermentation, the interferon containing E. coli cells are recovered from the broth for isolation and purification of rIFN- $\gamma$ . The following is a description of the fermentation and cell recovery processes.

##### 1. Preparation and Maintenance of Stock Cultures

A stock culture is prepared in sterile baffled culture flasks containing 150 to 500 mL of a sterile medium having the following composition.

Bactotryptone 10 g/L

Yeast Extract 5 g/L

Sodium Chloride 5-10 mg/L

The medium is then inoculated with a primary culture of E. coli W3110/py -CYC5.

The inoculated flask is then incubated on a shaker at 25-37°C until the absorbance at 550 nm reaches approximately 1.0. Approximately 50 percent v/v of 30 percent v/v dimethyl sulfoxide is added to the broth. One (1) mL aliquots are immediately dispensed into sterile vials and capped. The vials are stored at -60°C or below. Each fermentation is started using a replicate stock culture for inoculum.

##### 2. Inoculum Preparation

The inoculum is prepared in the medium previously described (L.B. Broth) in either shaker flasks or small fermenters. After incubation at about 37°C for approximately 8 hours, the inoculum is transferred to a fermenter. The volume of

the inoculum is between 2 to 10 percent of the volume of the fermentation.

### 3. Fermentation

Recombinant Interferon - Gamma production is carried out in fermenters with working volume of about 10 to 1000 liters.

The fermentation medium is composed of:

		<u>Per liter</u>
	*Glucose	50-100 g
	Ammonium Sulfate	4.0-8.0 g
	Potassium Phosphate, Monobasic	3.0-5.0 g
10	Potassium Phosphate, Dibasic	5.0-8.0 g
	Magnesium Sulfate, Heptahydrate	0.5-5.1 g
	Sodium Citrate, Dihydrate	0.5-2.0 g
	UCON LB-625	0.5-2.0 mL
	Ferric Chloride, Hexahydrate	0.005-0.15 g
	Zinc Sulfate, Heptahydrate	0.001-0.15 g
	Cobalt Chloride, Hexahydrate	0.001-0.005 g
15	Sodium Molybdate, Dihydrate	0.001-0.005 g
	Cupric Sulfate, Pentahydrate	0.001-0.005 g
	Boric Acid	0.001-0.005 g
	Manganese Sulfate, Monohydrate	0.001-0.005 g
	Hydrochloric Acid	0.0-1.0 mL
	Thiamine-HCl	0.0-0.1 g
	Tetracycline HCl	0.001-0.01 g
20	*L-Tryptophan	0.1-0.5 g
	Yeast Extract	2.0-8.0 g
	3- $\beta$ -Indoleacrylic acid	0.02-0.10 g

\*A portion of the glucose and tryptophan is added to the fermenter initially and the remainder is fed throughout the fermentation process.

Ingredients in the medium are sterilized by heat treatment or filtration prior to use in fermentation.

The fermentation is carried out at 25-40°C Other operating conditions are as follows:

	Agitation (rpm)	100-1000	
30	Aeration (vvm)	0.5-1.5	(Supplemented with oxygen when necessary)
	pH	6.5-7.5	(Controlled by the addition of ammonium hydroxide)

### 4. Purification

#### 35 a. Extraction of Recombinant Gamma Interferon.

E. coli cells are suspended in a medium which contains

salts and an appropriate buffer in the pH range of 6 to 9, preferably about 9. Recombinant gamma interferon is extracted by homogenization of the cell suspension in a high pressure colloid mill such as a Gaulin mill.

Sufficient polyethyleneimine is added to the solution to produce a 0.1 to 1% W/V solution. The supernatant contains gamma interferon.

b. Partial Purification of Recombinant Gamma Interferon on a Silica-based Adsorbant.

The supernatant from part (a) is adsorbed to a silica based adsorbant which is washed to remove impurities with appropriate salt solutions in the pH range of 6 to 9.

Recombinant gamma interferon is eluted using a solution containing 0.5 - 1.0 M tetramethyl ammonium chloride. All operations in this step are performed in the pH range of 7 to 9.

c. Partial Purification of Recombinant Gamma Interferon by Anion Exchange Chromatography

The eluent from part (b) is dialysed and adsorbed to an anion exchange chromatography medium which is then washed to remove impurities. Recombinant gamma interferon is eluted with a gradient of increasing salt. Typical anion exchange resins applicable for this step include carboxymethyl cellulose and sulphoethyl cellulose. All operations are performed in the pH range of between 7 and 9.

d. Partial Purification of Recombinant Gamma Interferon by Chromatography on Calcium Phosphate Gel

The eluent from part (c) is adsorbed to a medium of calcium phosphate which is then washed to remove impurities. The recombinant gamma interferon is eluted by increasing the salt concentration in a gradient of increasing phosphate concentration. All operations in this step are performed in the pH range of between 7 and 9.

e. Partial Purification of Recombinant Gamma Interferon by Anion Exchange Chromatography

The eluent from part (d) is dialysed and adsorbed to an anion exchange chromatography medium which is then washed to remove impurities. The recombinant gamma interferon is eluted from the anion exchange medium with a gradient of increasing salt concentration. Typical anion exchange chromatography media are carboxymethyl cellulose and sulphoethyl cellulose. All operations in this step are performed in the pH range of between 7 and 9.

f. Partial Purification of Recombinant Gamma Interferon by Gel Permeation Chromatography.

The eluent from part (e) is applied to a gel permeation medium and the column is developed with a salt containing medium. The appropriate recombinant gamma interferon containing fractions are pooled to produce the bulk drug substance. All operations in this step are performed in the pH range of between 7 and 9.

g. C-Terminal Amino Acid Sequence

To determine the c-terminal sequence samples were dialyzed into 70 percent formic acid, cleaved with cyanogen bromide and the resulting peptides separated on an Altex Ultrasphere C8 reverse phase HPLC column. Peaks were collected and analyzed by amino acid and sequence analysis. One C-terminal peptide was found:

-leu-phe-arg-gly-arg (residues 135-139, Figure 1). In some cases another additional peptide was detected:

-leu-phe-arg-gly-arg-arg-ala-ser-gln (residues 135-143, Figure 1). To determine the ratio of these two peptides, known amounts (by amino acid analysis) were loaded onto the reverse phase HPLC column and the respective peak heights determined. Each of three production lots contained less than about two percent of the long peptide (135-143, Figure 1), the balance being the 5-mer. This data is consistent with E. coli production of a mixture of 139 amino acid-containing and 143 amino acid-containing gamma interferons (excluding the N-terminal methionine, which is



also present in each case) in the relative proportions, respectively, of about 98.2 percent.

5. Formulation

Recombinant gamma interferon made in accordance with the foregoing is preferably formulated for parenteral administration according to the following Table.

Ingredient	QUANTITY PER VIAL	
	0.5 mg vial(1)	2.0 mg vial(2)
Recombinant Human Interferon - Gamma	0.5	2.0
Mannitol	100	80
Succinic Acid		
Disodium Hexahydrate	12.4	9.9
Glycine	5.6	4.5
Sodium Chloride	4.4	3.5
Polysorbate 20	0.8	0.6
Succinic Acid	0.5	0.4

- (1) Vials are reconstituted with 2.5 ml sterile Water for Injection.  
 (2) Vials are reconstituted with 2.0 ml sterile Water for Injection.

The interferons of the invention may be employed in medically appropriate dosage ranges, e.g., 1.0 mg/M<sup>2</sup> of body surface area.

D. Determination of the Activity of Various Gamma Interferons Following Trypsin Digestion

To establish the activity of various gamma interferons differing in their carboxy-termini gamma interferon prepared as described in the E. coli example, supra, was digested with trypsin to various degrees and tested by CPE assay with A549 cells as described within.

A sample of recombinant gamma interferon (r-HuIFN-gamma) (6.5 mg), prepared as described herein, was desalted over a small Sephadex G-25 molecular sieving column (PD-10, Pharmacia) into 0.10

M Ammonium Bicarbonate buffer, pH 8.5 to a final protein concentration of 2.1 mg/ml. A dilute trypsin solution (Worthington TPCK trypsin, 10 µg/ml in 0.001 M HCl, 16 µl) was added to 1.9 ml (4.0 mg) of the r-HuIFN-gamma solution, mixed and incubated at room temperature (trypsin:protein:1:25,000). Samples were removed from the incubation mixture at 1 hr., 3.5 hrs., 5.75 hrs., 8 hrs. and 10.25 hrs. At 8 hrs., an additional 15 µl (150 ng) of dilute trypsin solution was added in order to accelerate the reaction for the last time point sample at 10.25 hrs.

Fractionation of each time point sample into its respective components was performed on a Waters HPLC system using a BioRad Biogel HPHT Column. The time point aliquot in bicarbonate buffer was loaded directly (by manual injection) to the column at the time of sampling. The column was equilibrated in 0.01 M sodium phosphate pH 8.0, 30 µM calcium chloride and protein was eluted from the column utilizing a linear gradient of the equilibration buffer and 0.5 M sodium phosphate buffer pH 8.0, 0.6 µM calcium chloride.

Protein peaks as determined by absorbance at 214 nm and 280 nm were preparatively collected and stored covered at 4°C until analyzed. Typical analyses for selected peaks included:

1. Antiviral activity in the human lung carcinoma A549/EMC virus assay system (13).
2. SDS/PAGE fractionation by standard techniques employing the Laemmli gel system (14).
3. Protein concentration determination by the commercial (Pierce Chemical Co., Rockford, IL.) dye binding procedure.
4. Cyanogen bromide protein cleavage and subsequent HPLC analysis for peptide identification (15).

Samples of protein are dialyzed (12,000-14,000 mw cut off) overnight against 70% formic acid, taken to dryness by rotary evaporation and resuspended in 500 µl of 70% formic acid. Solid cyanogen bromide is added to each sample in a 12X75 mm glass tube, sealed, mixed until dissolved, covered with aluminum foil and incubated overnight at room temperature in a well ventilated area.

After cleavage, samples are taken to dryness by rotary

evaporation, resuspended in 0.5 ml of water and redried. Prior to fractionation by HPLC, samples were redissolved in 50% formic acid to a protein concentration of approximately 1 mg/ml.

Peptides were fractionated using a Waters HPLC system employing an Altex Ultrasphere Octyl column and a trifluoroacetic acid/water - trifluoroacetic acid/acetonitrile linear elution gradient. Wherever possible, peptides were identified by amino acid analysis. Table I lists comparative data for shortened forms of r-HuIFN- $\gamma$ :

10 Table 1

<u>r-HuIFN-<math>\gamma</math> Form*</u>	<u>C-Terminus**</u>	<u>Specific Activity (%)</u>
139aa:143aa::98:2 (for comparison)	LFRGR	100
15 131aa	AAKTGKRKR	40-50
129aa	AAKTGKR	6-9
125aa	AAK	ca. 1

\* Based on numbering system of Figure 1, but excluding the N-terminal methionine also present in each case.

20 \*\* Conventional single letter abbreviations for amino acid residues:

25 A = alanine  
F = phenylalanine  
G = glycine  
K = lysine  
L = leucine  
R = arginine  
T = threonine

30 It will be appreciated that gamma interferons of any length within the range 126aa-143aa (excluding N-terminal methionine) will be expressed from appropriately tailored genes. Thus, for example, the gene depicted in Figure 1 contains a Fnu4H restriction site at

123 124  
Ala Ala

GCA GCT  
CGT CGA

35 Following restriction with Fnu4H, synthetic oligonucleotides

encoding any desired sequence followed by a "stop" codon and a linker compatible with an available restriction site in the expression plasmid may be ligated to the fore part of the gamma interferon gene. For example, the sequence

5

A GCT AAA ACA      TA  
                           -(X)-  
       CGA TTT TGT      ATCTAG

wherein X encodes one or more amino acids may be ligated into the E. coli expression vehicle exemplified above following digestion of the plasmid with Fnu4H and BglII, a stop codon resulting from ligation thusly

10

                          stop  
                           |  
                   ...(X)-TAGATC...  
                           |  
                   ATCTAG  
                           BglII

15

#### E. Assay: Inhibition of Cytopathic Effect (CPE)

##### 1. Test Procedure

20

To each well add 100 microliters of a suspension of human lung carcinoma (A549) cells (ATCC No. CCL 185) which have been adjusted to contain  $4 \times 10^5$  cells/ml in Eagles MEM.

Incubate plates at 37°C for approximately 18 hours.

25

After 18 to 24 hours incubation, add to each well in the first column, 80 microliters additional medium.

Add to a well in the first column 20 microliters of a sample to be assayed for interferon activity.

30

Transfer 100 microliters of the contents of each well in the first column horizontally to each well in second column.

35

Continue to transfer 100 microliters of the contents of a well from column to subsequent column until a total of 10 transfers are performed ending in column 11.

After 24 hours of incubation, challenge all wells, except cell controls, with 50 microliters of encephalomyocarditis virus at a multiplicity of infection which results in 100% cytopathic effect in 24 hours after infection.

5

Cover trays with lids and incubate at 37°C for 24 hours.

Pour off fluid from all wells and stain 5-15 minutes with 0.5% crystal violet.

10

Viability of cells is determined by observation of stained cells.

Titer of a sample is the reciprocal of the dilution where 50% viable cells remain.

15

## 2. Calculation

The activity of all samples is normalized by the Reference Units Conversion Factor which is calculated from:

20

$$\frac{\text{Actual Titer of Standard Interferon}}{\text{Observed Titer}} = \frac{\text{Reference Units}}{\text{Conversion Factor}} = \text{RUCF}$$

## 3. Specific Activity

Using the A549/EMCV bioassay systems standardized with IFN-gamma reference material provided by the NIH, the specific antiviral activity of recombinant human IFN-gamma is approximately three times higher than the activity of the modified rIFN-gamma molecule with three additional amino acids (cys-tyr-cys) at the N-terminus.

25

## 4. Stability

Based again on the above mentioned measure of bioactivity (A549/EMCV), formulated and vialled rIFN-gamma continues to be stable (no loss of biological activity) three months after production (stored at 4°C).

30

## F. Antiproliferative activity of rIFN-gamma compared to the other types of human interferon

35

## 1. Materials and Methods

- 5           rIFN-gamma:       Des Cys-Tyr-Cys recombinant interferon gamma in  
                          the form of a solution (20 mM sodium succinate,  
                          0.15 M NaCl, pH 6). The material was prepared  
                          in accordance with the E. coli Example above.  
                          The specific activity was  $2.7 \times 10^7$  IU/mg  
                          protein.
- 10          CTC-rIFN-gamma: Recombinant interferon gamma having  
                          "Cys-Tyr-Cys" structure at the N-terminus of  
                          the molecule in the form of a solution (20 mM  
                          sodium succinate, 0.15 M NaCl, pH 6). The  
                          specific activity was  $1.3 \times 10^7$  IU/mg protein.
- 15          HuIFN-beta:    Lyophilized human fibroblast interferon  
                          produced in human diploid foreskin fibroblast  
                          with a specific activity of more than  $1 \times 10^7$   
                          IU/mg protein prepared by Toray Ind., Inc. A  
                          vial contains  $3 \times 10^6$  IU HuIFN-beta and 3 mg  
                          human serum albumin.
- 20          HuIFN-alpha:   Human natural leukocyte interferon with a  
                          specific activity of  $4 \times 10^6$  IU/mg protein  
                          supplied in the form of solution by Dr. K.  
                          Cantell, Central Public Health Laboratory,  
                          Helsinki, Finland.
- 25          Control:        Placebo containing 3 mg human serum albumin.
- 30          Culture Medium: Eagle's minimum essential medium supplemented  
                          with 10 percent heat-inactivated precolostrum  
                          newborn calf serum ("PNCS") and 2 mM  
                          L-glutamine was used with Hela, KB, HMV-1, FL,  
                          and J-111 cells. Dulbecco's minimum essential  
                          medium containing 10 percent heat-inactivated  
                          PNCS, 100  $\mu$ g/ml kanamycin, and 2 mM L-glutamine  
                          was used with A549 cells. RPMI 1640 medium  
                          supplemented with heat-inactivated PNCS and  
                          100  $\mu$ g/ml kanamycin was used with the remaining  
35                           human cells listed in Table 2.

## 2. Evaluation of antiproliferative activity

The test cells suspended in the culture medium were seeded into plastic tissue culture plates at a concentration of  $5 \times 10^3$  cells/0.5 ml/well. Various amounts of interferon dissolved in the corresponding culture medium (0.5 ml) were added subsequently (day 0). Cultivation was carried out at 37°C in a humidified atmosphere of 5 percent CO<sub>2</sub> and 95 percent air. On day 6 the culture media were removed and the cells in suspension culture were directly suspended in Isoton II (Coulter Electronics Inc.) for cell counting in a Coulter counter. The cells forming a sheet in plastic vessels were pre-treated with 0.05 percent trypsin-0.02 percent EDTA to prepared single cell suspension in Isoton II (Coulter Electronics Inc.). Antiproliferative activity of interferon was expressed as the antiviral units required to produce 50 percent reduction of cell number (IC<sub>50</sub>, IU/ml) compared to the control culture (without interferon).

As shown in the table, the antiproliferation activity of rIFN-gamma varied markedly depending on the human cell species. In this case KATO-III, siglet-ring cell carcinoma of the stomach, was highly sensitive, the cell line showing the IC<sub>50</sub> of 1.2 IU/ml, while Daudi cells, Burkitt's lymphoma, which were highly sensitive to type II interferon (HuIFN-alpha, HuIFN-beta), were insensitive to type II interferon including rIFN-gamma. Lung adenocarcinoma (PC-8, PC-12) was insensitive to all interferon species tested. The anticellular spectrum between rIFN-gamma and CTC-rIFN-gamma was almost the same and it was generally apparent that the antiproliferative efficacy of rIFN-gamma was superior to that of CTC-rIFN-gamma. In the case of comparison between four interferons, the highest efficacy was obtained from rIFN-gamma except for Daudi cells.

### G. Comparison of stability between rIFN-gamma and CTC-rIFN-gamma in various fluids in vitro

$1 \times 10^6$  IU/vial of lyophilized rIFN-gamma prepared in accordance with the E. coli Example above and  $1 \times 10^6$  IU/vial of lyophilized rIFN-gamma having a "Cys-Tyr-Cys" structure at the

TABLE 2

ANTIPROLIFERATIVE ACTIVITY OF rIFN- $\gamma$  COMPARED TO OTHER INTERFERON SPECIES

Cell	Histological Type or Origin of Test Cells	IC50 (IU/ml)			
		rIFN- $\gamma$	CTC-rIFN- $\gamma$	HuIFN- $\beta$	HuIFN- $\alpha$
KATO-III	Siglet-ring cell carcinoma of stomach	1.2	1.3	--	--
MKN-1	Adenosquamous carcinoma of stomach	37	330	--	--
MKN-28	Well differentiated adenocarcinoma of stomach	$>10^3$	$>10^3$	--	--
MKN-45	Poorly differentiated adenocarcinoma of stomach	$>10^3$	$>10^3$	--	--
MKN-74	Well differentiated adenocarcinoma of stomach	$>10^3$	$>10^3$	--	--
HeLa	Uterine cervix carcinoma	22	132	446	$>10^3$
KB	Nasopharyngeal carcinoma	54	330	$>10^4$	$>10^4$
HMV-1	Amelanotic melanoma	38	45	130	820
SEKI-F	Melanoma	135	$>10^3$	137	663
FL	Amnion (non-malignant origin)	38	45	$>10^3$	$>10^4$
PC-8	Lung poorly differentiated adenocarcinoma	$>10^3$	$>10^3$	$>10^4$	$>10^3$
PC-12	Lung adenocarcinoma	$>10^3$	$>10^3$	$>10^4$	$>10^3$
A549	Lung alveolar carcinoma	55	--	--	--
QG56	Lung squamous cell carcinoma	5	--	--	--
QG90	Lung anaplastic small cell carcinoma	5.5	--	--	--
Daudi	Burkitt's lymphoma	$>10^3$	$>10^3$	47	7
Namaiva	Burkitt's lymphoma	$>10^3$	$>10^3$	--	--
J-111	Monocytic leukemia	25	84	204	$>10^3$



N-terminus, with 10 mg human serum albumin, phosphate buffer and an isotonic amount of NaCl for each interferon were reconstituted with distilled water and the concentration was adjusted to  $4 \times 10^4$  IU/ml.

5        In vitro stability of interferons was evaluated by the determination of the residual anti-viral activity in various fluids. Incubations were initiated by addition of the above interferon solutions into nine volumes of rabbit serum, human serum or Eagle's MEM which were pre-incubated 10 min. in a water bath incubator at 37°C or 4°C. At 0, 0.25, 0.5, 1, 4, 8, 24, 72 and 10        144 hr., an aliquot was harvested and mixed with the nine volumes of Eagle's MEM. The samples were kept frozen at -80°C in a deep freezer until the assay of interferon titer. Interferon titer was assayed by the CPE50 reduction method using human amniotic cells (FL 15        cell) challenged with Sindbis virus. The results are indicated in Figure 3 as percentages of residual titer against the additional titer.

While the invention has been exemplified by reference to the preferred embodiments in which expression is effected in E. coli and 20        in COS-7 cells, it will be apparent that the recombinant gamma interferons of the invention may be produced in other systems as well such as other bacterial strains, yeast and tissue culture systems, as to which see GB 2107718 and (6). Thus, the invention is not confined to the most preferred 25        embodiments, but instead extends to all legally cognizable equivalents of the claims which follow.

30

35

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CLAIMS:

1. A composition suitable for pharmaceutical administration including

- 5 a) a recombinant gamma interferon polypeptide comprising the amino acid sequence, extending from the N-terminus:

X-Y-ASP-PRO-TYR-VAL-LYS-GLU-ALA-GLU-ASN-LEU-LYS-LYS-TYR-PHE-  
 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15  
 10 ASN-ALA-GLY-HIS-SER-ASP-VAL-ALA-ASP-ASN-GLY-THR-LEU-PHE-LEU-  
 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30  
 GLY-ILE-LEU-LYS-ASN-TRP-LYS-GLU-GLU-SER-ASP-ARG-LYS-ILE-MET-  
 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45  
 GLN-SER-GLN-ILE-VAL-SER-PHE-TYR-PHE-LYS-LEU-PHE-LYS-ASN-PHE-  
 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60  
 15 LYS-ASP-ASP-GLN-SER-ILE-GLN-LYS-SER-VAL-GLU-THR-ILE-LYS-GLU-  
 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75  
 ASP-MET-ASN-VAL-LYS-PHE-PHE-ASN-SER-ASN-LYS-LYS-LYS-ARG-ASP-  
 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90  
 20 ASP-PHE-GLU-LYS-LEU-THR-ASN-TYR-SER-VAL-THR-ASP-LEU-ASN-VAL-  
 91 92 93 94 95 96 97 98 99 100 101 102 103 104 105  
 GLN-ARG-LYS-ALA-ILE-HIS-GLU-LEU-ILE-GLN-VAL-MET-ALA-GLU-LEU-  
 106 107 108 109 110 111 112 113 114 115 116 117 118 119 120  
 SER-PRO-ALA-ALA-LYS-THR  
 121 122 123 124 125 126

25 wherein X is a methionine residue or hydrogen and Y is a glutamine residue or, where X is hydrogen, either a glutamine or a pyroglutamate residue; or

- b) a modified species of gamma interferon polypeptide of a) having equivalent properties thereof.

30

2. A composition according to claim 1 wherein X is methionine.

3. A composition according to claim 2 wherein said polypeptide has 126+n amino acids extending beyond X, n being zero or an integer from 1 to 17.

35

4. A composition according to claim 3 wherein the n amino acids are

GLY-LYS-ARG-LYS-ARG-SER-GLN-MET-LEU-  
127 128 129 130 131 132 133 134 135

PHE-ARG-GLY-ARG-ARG-ALA-SER-GLN  
136 137 138 139 140 141 142 143

5

5. A composition according to claim 3 wherein the polypeptide is selected from analogs of the polypeptide

in which amino acid 143 is absent or in which the amino acid sequence Z to amino acid 143 is absent, Z being amino acid 127, 128, 129, 130, 131, 132, 133, 134, 135, 136, 137, 138, 139, 140, 141 or 142.

10

6. A composition according to any of claims 1 to 5 whose polypeptide exhibits activity in cytopathic inhibition microtiter assay at least about equal to that of a specified recombinant gamma interferon whose amino acid sequence is

15

X-A-GLN-ASP-PRO-TYR-VAL-LYS-GLU-ALA-GLU-ASN-LEU-LYS-LYS-TYR-PHE-ASN-  
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16

20

ALA-GLY-HIS-SER-ASP-VAL-ALA-ASP-ASN-GLY-THR-LEU-PHE-LEU-GLY-ILE-LEU-  
17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33

LYS-ASN-TRP-LYS-GLU-GLU-SER-ASP-ARG-LYS-ILE-MET-GLN-SER-GLN-ILE-VAL-  
34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50

SER-PHE-TYR-PHE-LYS-LEU-PHE-LYS-ASN-PHE-LYS-ASP-ASP-GLN-SER-ILE-GLN-  
51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67

25

LYS-SER-VAL-GLU-THR-ILE-LYS-GLU-ASP-MET-ASN-VAL-LYS-PHE-PHE-ASN-SER-  
68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84

ASN-LYS-LYS-LYS-ARG-ASP-ASP-PHE-GLU-LYS-LEU-THR-ASN-TYR-SER-VAL-THR-  
85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100 101

30

ASP-LEU-ASN-VAL-GLN-ARG-LYS-ALA-ILE-HIS-GLU-LEU-ILE-GLN-VAL-MET-ALA-1  
02 103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118

GLU-LEU-SER-PRO-ALA-ALA-LYS-THR-GLY-LYS-ARG-LYS-ARG-SER-GLN-MET-LEU-1  
19 120 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135

PHE-ARG-GLY-ARG-ARG-ALA-SER-GLN  
136 137 138 139 140 141 142 143

35

wherein X is hydrogen and A is CYS-TYR-CYS.

7. A composition according to claim 6  
exhibiting activity in cytopathic effect inhibition  
microtiter assay at least about twice that of said  
specified recombinant gamma interferon.
- 5
8. A replicable expression vehicle capable, in a trans-  
formant microorganism or cell culture, of expressing a  
recombinant gamma interferon polypeptide as defined in  
any one of claims 1 to 5.
- 10
9. An expression vehicle according to claim 8 for use in a  
microorganism.
10. An expression vehicle according to claim 8 for use in a  
15 bacterium.
11. An expression vehicle according to any one of claims 8,  
9 and 10 wherein the expression of said polypeptide is  
unaccompanied by the formation of any associated signal  
20 peptide.
12. A transformed microorganism or cell comprising a host  
microorganism or cell which has been transformed with  
the replicable expression vehicle of any of claims 8 to  
25 11.
13. A double-stranded DNA comprising a coding strand and a  
complementary strand wherein the coding strand encodes a  
polypeptide comprising the amino acid sequence, extending from  
30 the N-terminus of said polypeptide,
- MET-GLN-ASP-PRO-TYR-VAL-LYS-GLU-ALA-GLU-ASN-LEU-LYS-LYS-TYR-PHE-  
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15
- ASN-ALA-GLY-HIS-SER-ASP-VAL-ALA-ASP-ASN-GLY-THR-LEU-PHE-LEU-  
16 17 18 19 20 21 22 23 24 25 26 27 28 29 30
- 35 GLY-ILE-LEU-LYS-ASN-TRP-LYS-GLU-GLU-SER-ASP-ARG-LYS-ILE-MET-  
31 32 33 34 35 36 37 38 39 40 41 42 43 44 45
- GLN-SER-GLN-ILE-VAL-SER-PHE-TYR-PHE-LYS-LEU-PHE-LYS-ASN-PHE-  
46 47 48 49 50 51 52 53 54 55 56 57 58 59 60

LYS-ASP-ASP-GLN-SER-ILE-GLN-LYS-SER-VAL-GLU-THR-ILE-LYS-GLU-  
61 62 63 64 65 66 67 68 69 70 71 72 73 74 75

ASP-MET-ASN-VAL-LYS-PHE-PHE-ASN-SER-ASN-LYS-LYS-LYS-ARG-ASP-  
76 77 78 79 80 81 82 83 84 85 86 87 88 89 90

5 ASP-PHE-GLU-LYS-LEU-THR-ASN-TYR-SER-VAL-THR-ASP-LEU-ASN-VAL-  
91 92 93 94 95 96 97 98 99 100 101 102 103 104 105

GLN-ARG-LYS-ALA-ILE-HIS-GLU-LEU-ILE-GLN-VAL-MET-ALA-GLU-LEU-  
106 107 108 109 110 111 112 113 114 115 116 117 118 119 120

10 SER-PRO-ALA-ALA-LYS-THR  
121 122 123 124 125 126.

14. A double stranded DNA according to claim 13 wherein  
said polypeptide from amino acid no. 126 is as defined  
in any one of claims 3, 4 and 5.

15 15. A polypeptide exhibiting the characteristic properties  
of native human gamma interferon and containing at  
least the amino acid sequence set forth in claim 1.

20 16. A polypeptide of claim 15, wherein the amino acid  
sequence extends beyond position 126 and is:

GLY-LYS-ARG-LYS-ARG-SER-GLN-MET-LEU-PHE-ARG-GLY-ARG  
127 128 129 130 131 132 133 134 135 136 137 138 139

25 17. A polypeptide of claim 15, wherein the amino acid sequence  
extends beyond position 126 and is:

GLY-LYS-ARG-LYS-ARG-SER-GLN-MET-LEU-PHE-ARG-GLY-ARG-ARG-ALA-SER-GLN  
127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143

30 18. A polypeptide of Claim 15, wherein the amino acid sequence extends  
beyond position 126 and is:

GLY-LYS-ARG  
127 128 129

35 19. A polypeptide of Claim 15, wherein the amino acid sequence extends  
beyond position 126 and is:

GLY-LYS-ARG-LYS-ARG  
127 128 129 130 131

20. A polypeptide of Claim 15, wherein the amino acid sequence extends beyond position 126 and is:

GLY-LYS-ARG-LYS-ARG-Z  
127 128 129 130 131

wherein Z is H,

5 SER  
132,  
SER-GLN  
132 133,  
SER-GLN-MET  
132 133 134,  
SER-GLN-MET-LEU  
10 132 133 134 135,  
SER-GLN-MET-LEU-PHE  
132 133 134 135 136,  
SER-GLN-MET-LEU-PHE-ARG  
132 133 134 135 136 137,  
SER-GLN-MET-LEU-PHE-ARG-GLY  
132 133 134 135 136 137 138, or  
15 SER-GLN-MET-LEU-PHE-ARG-GLY-ARG  
132 133 134 135 136 137 138 139.

21. A polypeptide of Claim 15, wherein the amino acid sequence extends beyond position 126 and is:

GLY-LYS-ARG-LYS-ARG-Z  
127 128 129 130 131

20

wherein Z is H,

SER  
132,  
SER-GLN  
132 133,  
SER-GLN-MET  
25 132 133 134,  
SER-GLN-MET-LEU  
132 133 134 135,  
SER-GLN-MET-LEU-PHE  
132 133 134 135 136,  
SER-GLN-MET-LEU-PHE-ARG  
132 133 134 135 136 137,  
SER-GLN-MET-LEU-PHE-ARG-GLY  
30 132 133 134 135 136 137 138,  
SER-GLN-MET-LEU-PHE-ARG-GLY-ARG  
132 133 134 135 136 137 138 139,  
SER-GLN-MET-LEU-PHE-ARG-GLY-ARG-ARG  
132 133 134 135 136 137 138 139 140,  
SER-GLN-MET-LEU-PHE-ARG-GLY-ARG-ARG-ALA  
132 133 134 135 136 137 138 139 140 141,  
35 SER-GLN-MET-LEU-PHE-ARG-GLY-ARG-ARG-ALA-SER  
132 133 134 135 136 137 138 139 140 141 142, or  
SER-GLN-MET-LEU-PHE-ARG-GLY-ARG-ARG-ALA-SER-GLN  
132 133 134 135 136 137 138 139 140 141 142 143.

22. A composition comprising a mixture of the polypeptides of Claim 16 and 17.
- 5 23. A composition of Claim 22 comprising greater than about 95% of the polypeptide of Claim 18.
24. A composition of Claim 22 comprising greater than about 97% of the polypeptide of Claim 18.
- 10 25. A composition according to any one of claims 1 to 7 comprising a pharmaceutically acceptable level of human serum albumin.



14

TGAAGATCAGCTATTAGAAGAGAAAGATCAGTTAAGTCCTTTGGACCTGATCAGCTTGATACAAG  
50

s1  
met lys tyr thr ser  
**AACTACTGATTCAACTTCTTTGGCTTAATTCTCTCGGAAACG** ATG AAA TAT ACA AGT  
100

tyr ile leu ala phe gln leu cys ile val leu gly ser leu  
 TAT ATC TTG GCT TTT CAG CTC TGC ATC GTT TTG GGT TCT CTT

gly cys tyr cys s23 1 10  
GCG TGT TAC TGC CAG GAC CCA TAT GTA AAA GAA GCA GAA AAC CTT AAG

20  
 LYS TYR PHE ASN ALA GLY HIS SER ASP VAL ALA ASP ASN GLY  
 AAA TAT TTT AAT GCA GGT CAT TCA GAT GTA GCG GAT AAT GGA  
 250

30  
 THR LEU PHE LEU GLY ILE LEU LYS ASN TRP LYS GLU GLU SER ASP ARG  
 ACT CTT TTC TTA GGC ATT TTG AAG AAT TGG AAA GAG GAG AGT GAC AGA  
 300

50  
 LYS ILE MET GLN SER GLN ILE VAL SER PHE TYR PHE LYS LEU  
 AAA ATA ATG CAG AGC CAA ATT GTC TCC TTT TAC TTC AAA CTT

60
70  
PHE LYS ASN PHE LYS ASP ASP GLN SER ILE GLN LYS SER VAL GLU THR  
TTT AAA AAC TTT AAA GAT GAC CAG AGC ATC CAA AAG AGT GTG GAG ACC  
350

80  
ILE LYS GLU ASP MET ASN VAL LYS PHE PHE ASN SER ASN LYS  
ATC AAG GAA GAC ATG AAT GTC AAG TTT TTC AAT AGC AAC AAA  
400

90 110  
 LYS LYS ARG ASP ASP PHE GLU LYS LEU THR ASN TYR SER VAL THR ASP  
 AAG AAA CGA GAT GAC TTC GAA AAG CTG ACT AAT TAT TCG GTA ACT GAC  
 450

LEU ASN VAL GLN ARG LYS ALA ILE HIS GLU LEU ILE GLN VAL  
TTG AAT GTC CAA CGC AAA GCA ATA CAT GAA CTC ATC CAA GTG

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Fig. 1. (Cont.)

120 130  
MET ALA GLU LEU SER PRO ALA ALA LYS THR GLY LYS ARG LYS ARG SER  
ATG GCT GAA CTG TCG CCA GCA GCT AAA ACA GGG AAG CGA AAA AGG AGT  
550

140 143 STOP  
GLN MET LEU PHE ARG GLY ARG ARG ALA SER GLN  
CAG ATG CTG TTT CGA GGT CGA AGA GCA TCC CAG TAA TGGTTGT  
600

CCTGCCTGCAATATTTGAATTTTAAATCTAAATCTATTTATTAATATTTAACATTATTTATATG  
650

GGGAATATATTTTAGACTCTCATCAATCAAATAAGTATTTATAATAGCAACTTT  
700

TGTGTAATGAAAATGAATATCTATTAATATATGTATTATTTATAATTCCTATATCCTGTGACTG  
750

TCTCACTTAATCCTTTGTTTTCTGACTAATTAGGCAAGGCTATGTGATTACAAGG  
800 850

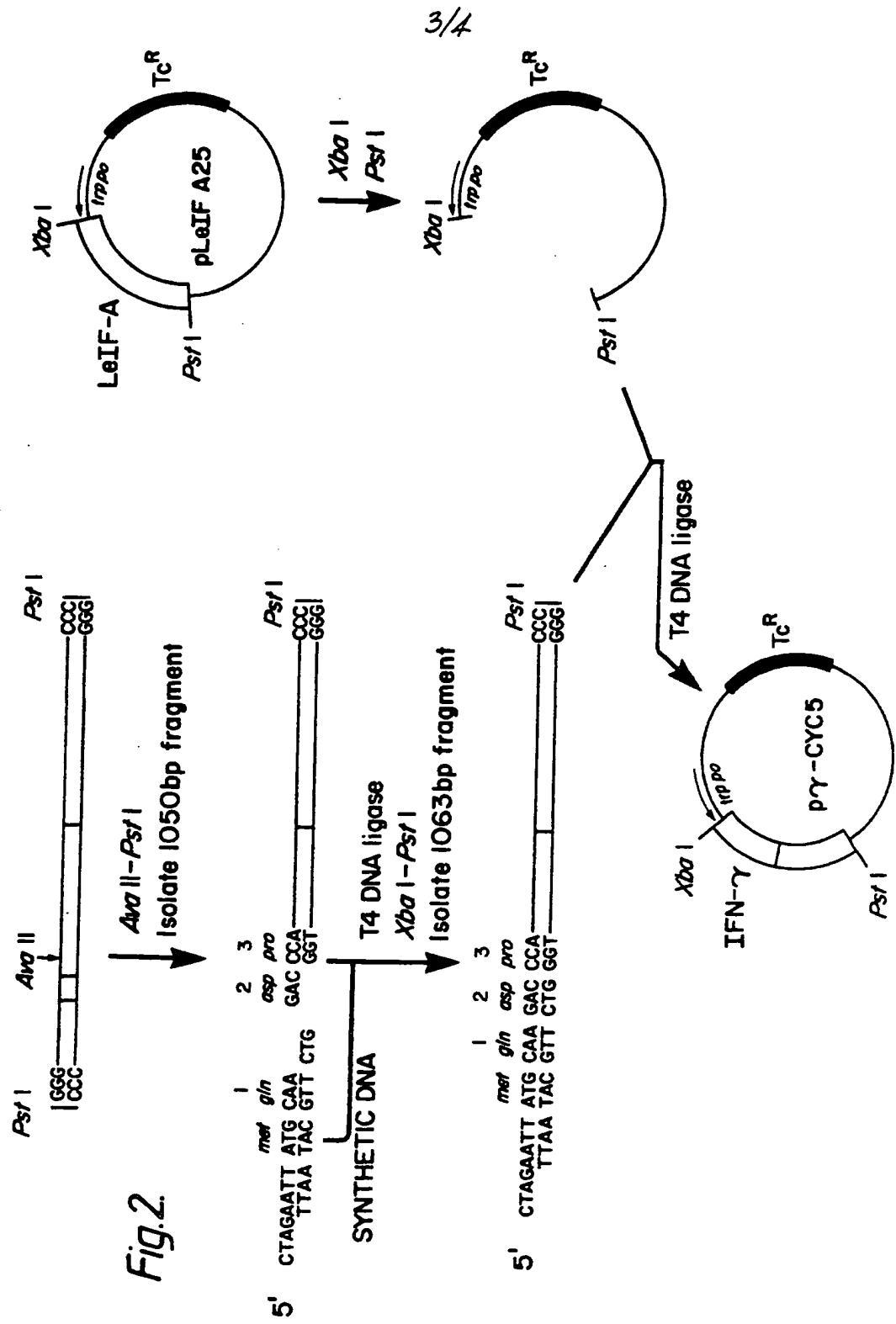
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900

ACTTGATGATACAATGAACACTTATAAGTGAAGTGATACTATCCAGTTACTGCCG  
950

GTTTGAAAATATGCCTGCAATCTGAGCCAGTGCTTTAATGGCATGTCAGACAGAAGTTGAATGT  
1000

GTCAGGTGACCCTGATGAAAACATAGCATCTCAGGAGATTTTCATGCCTGGTGCTT  
1050

CCAAATATTGTTGACAACTGTGACTGTACCCAAATGGAAAGTAACTCATTGTTAAAATTATCA  
1100 1150



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Fig. 3.

